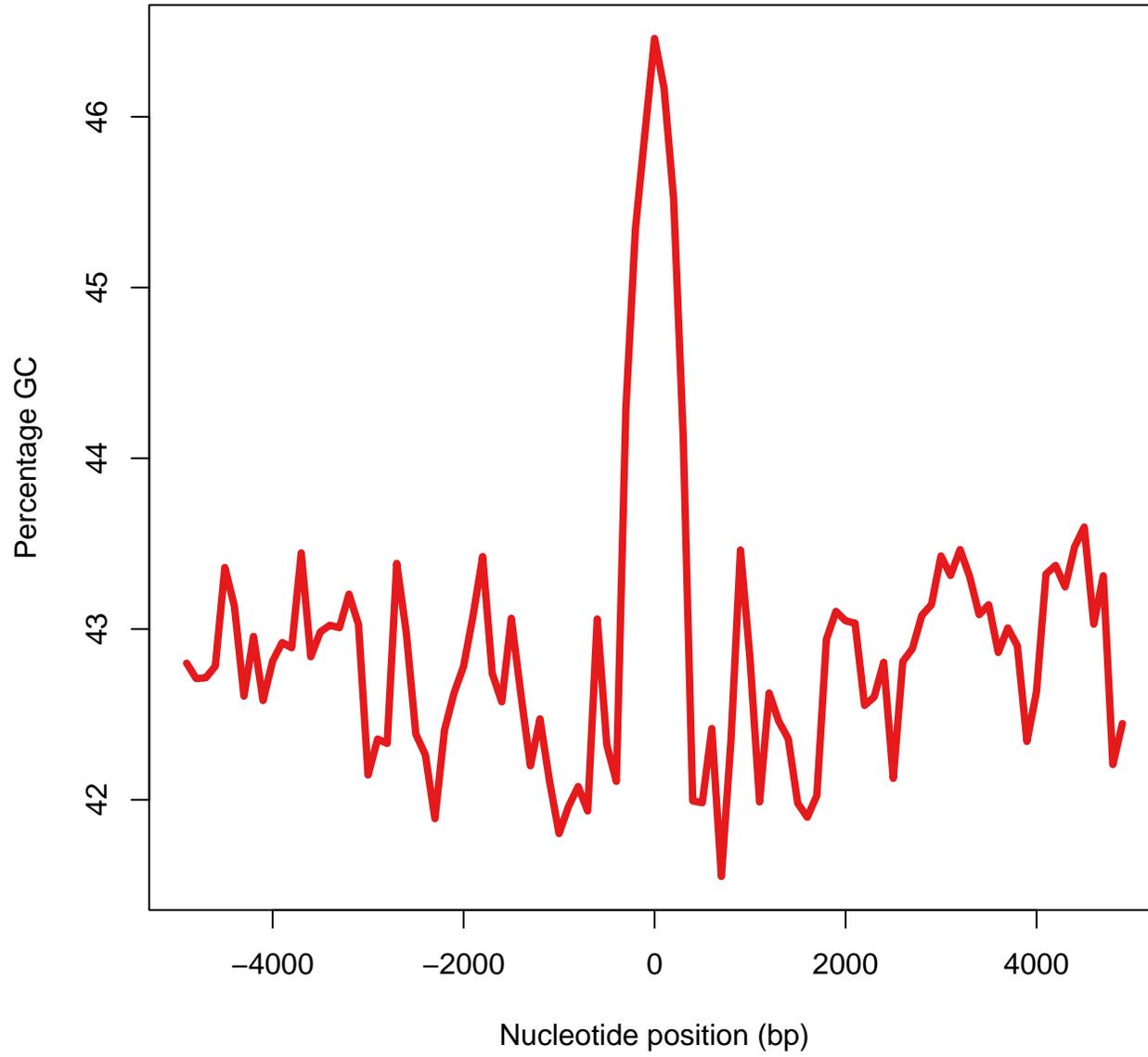
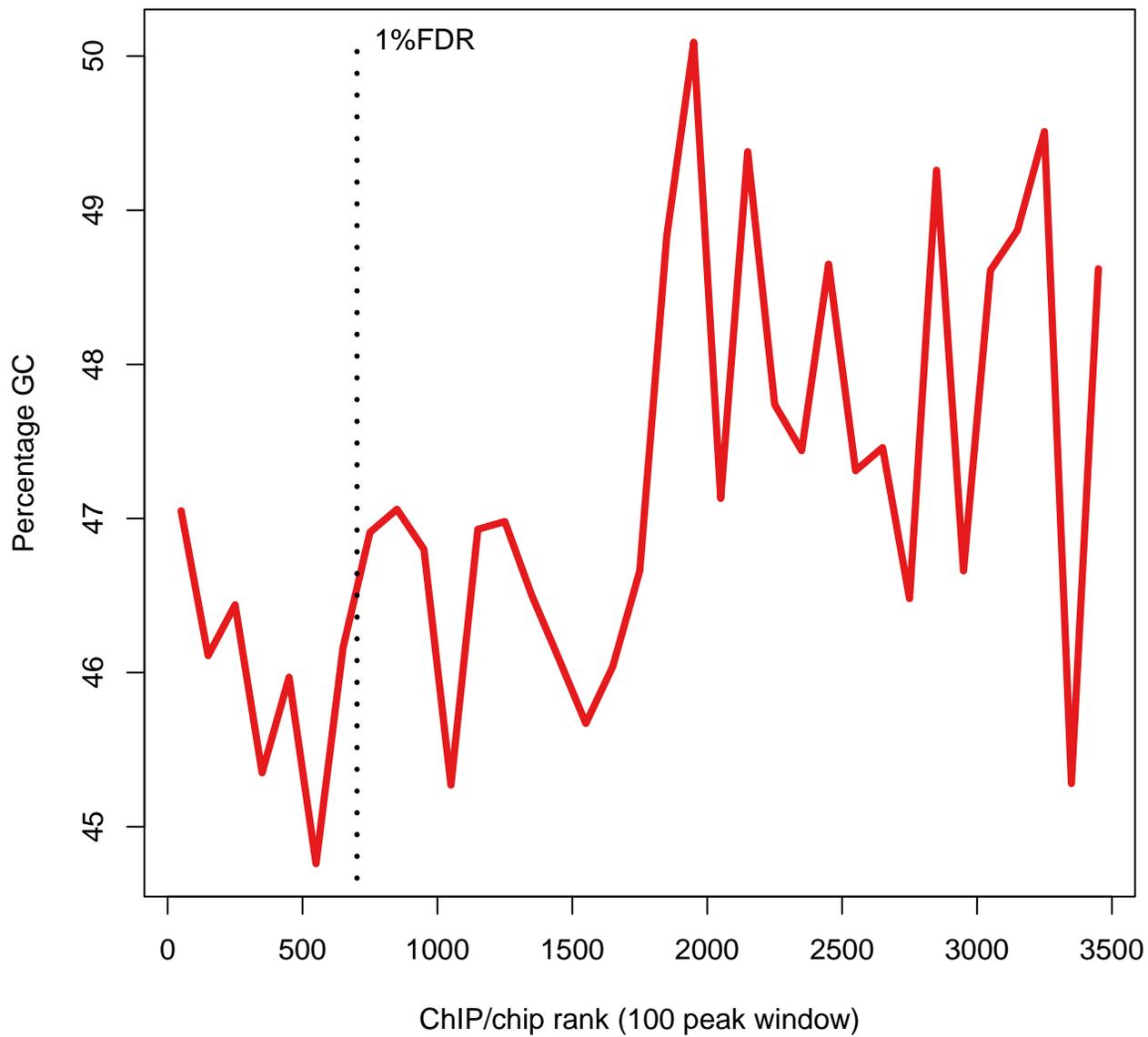


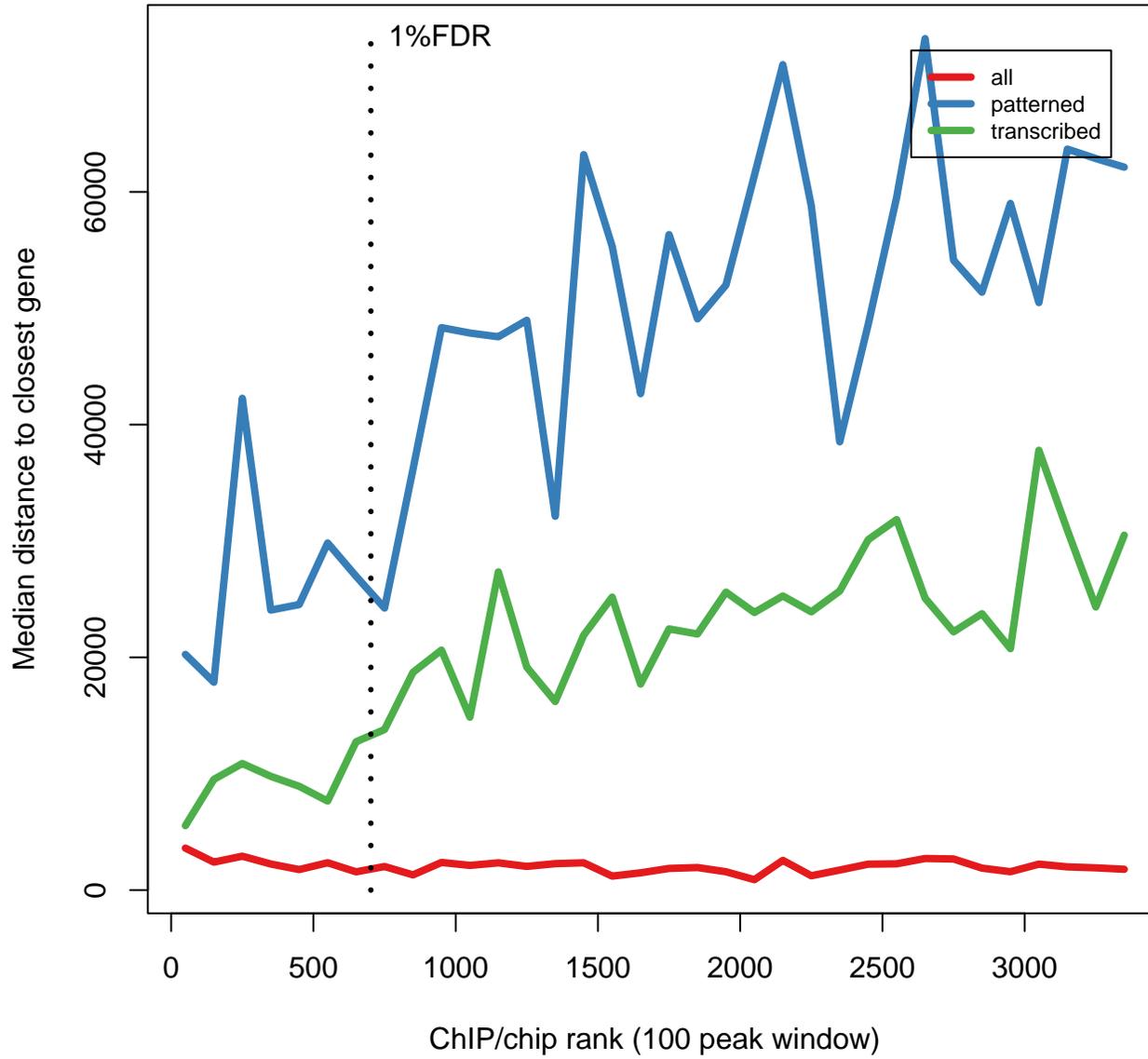
**bcd\_2 base composition across peaks**



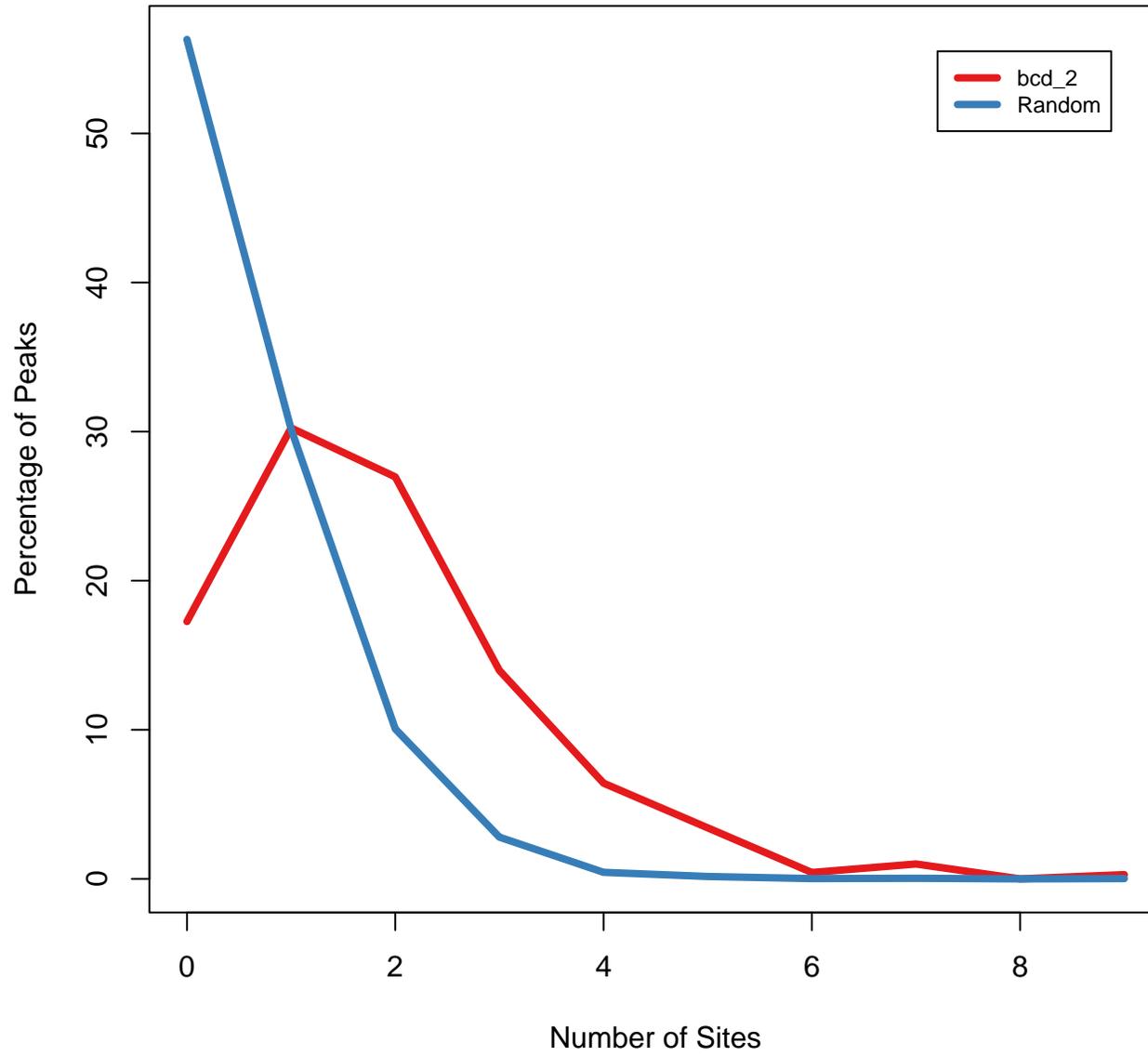
### bcd\_2 base composition down ranks



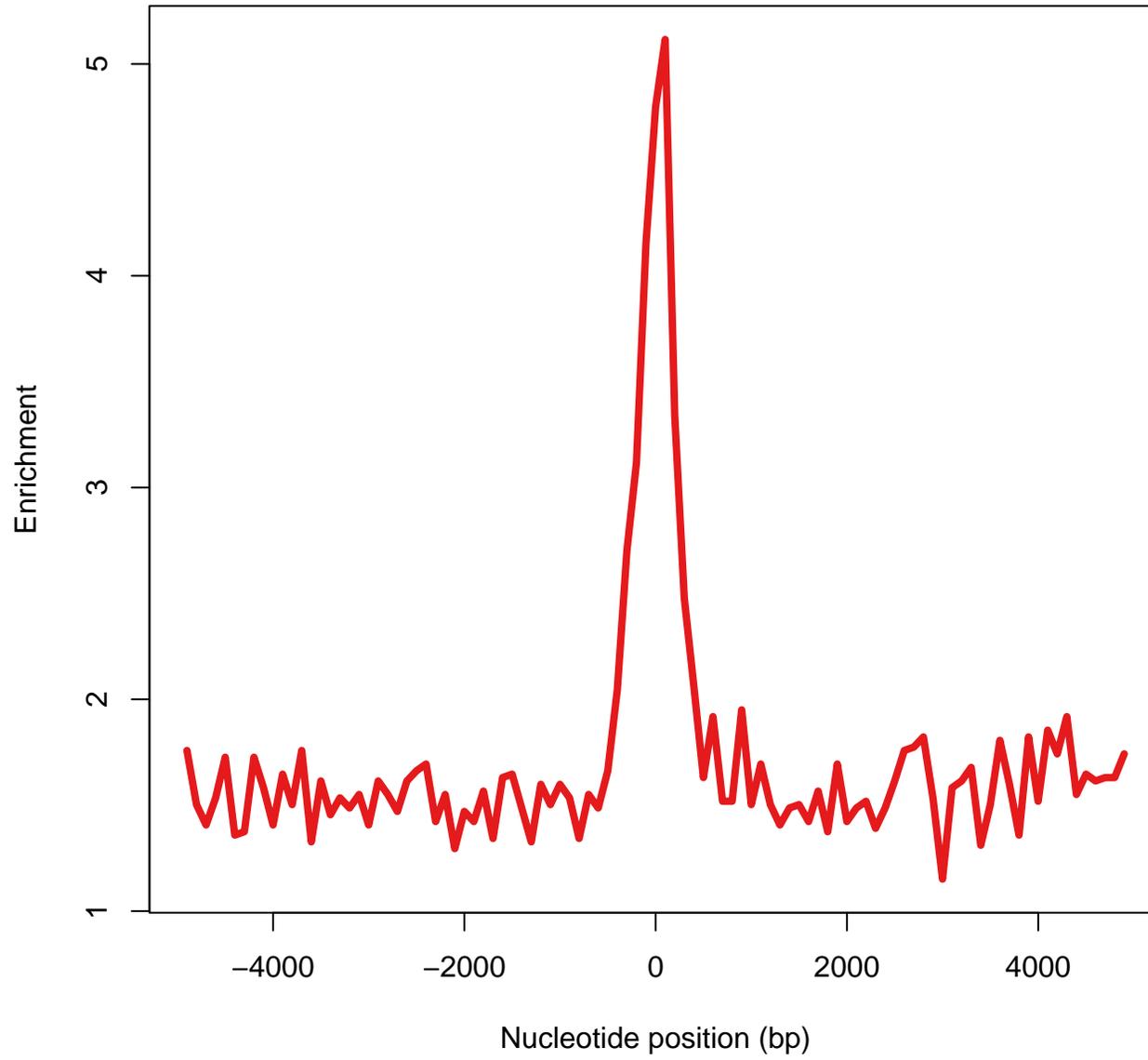
bcd\_2 median distance to genes



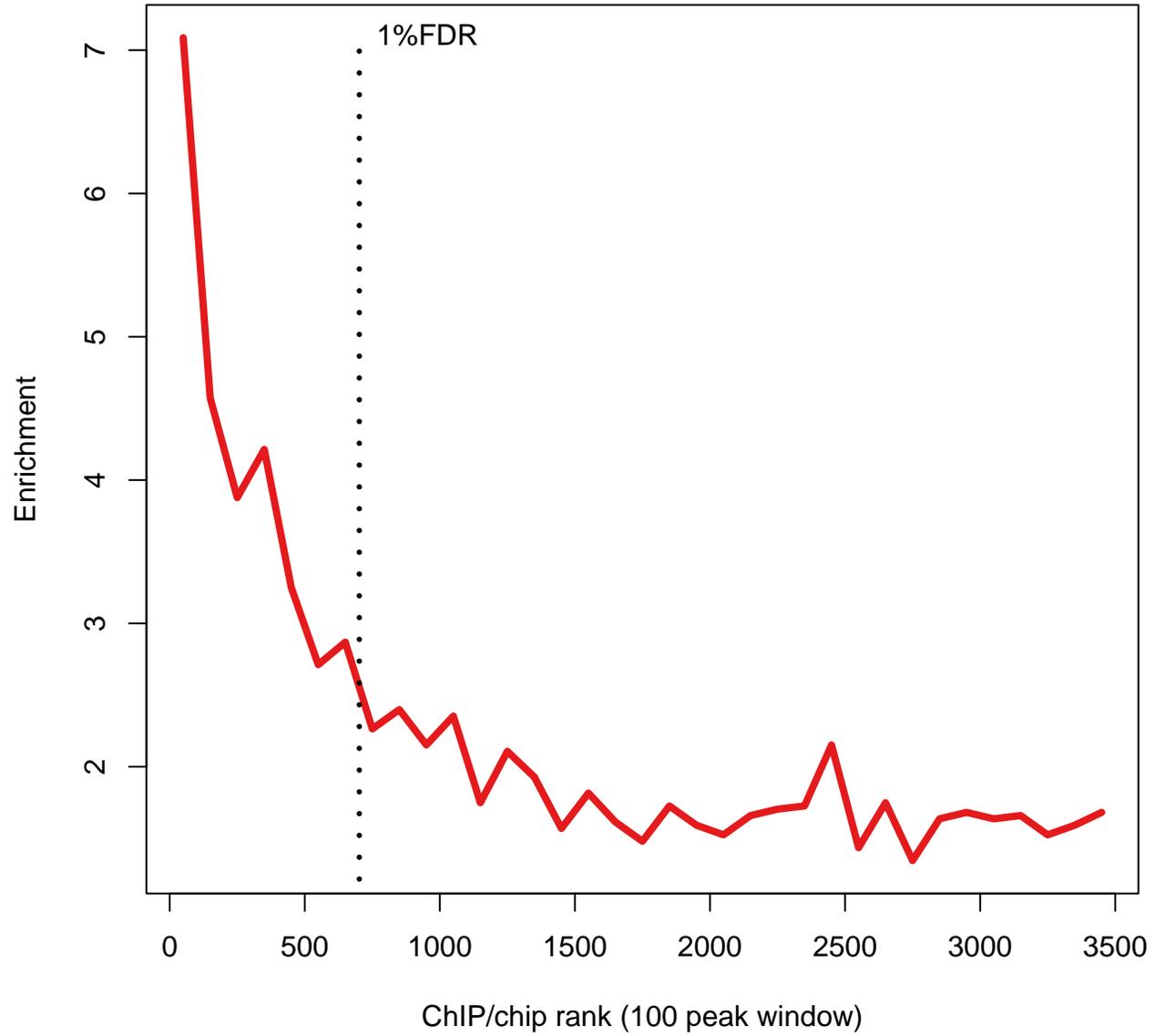
**bcd\_2 distribution of number of sites**



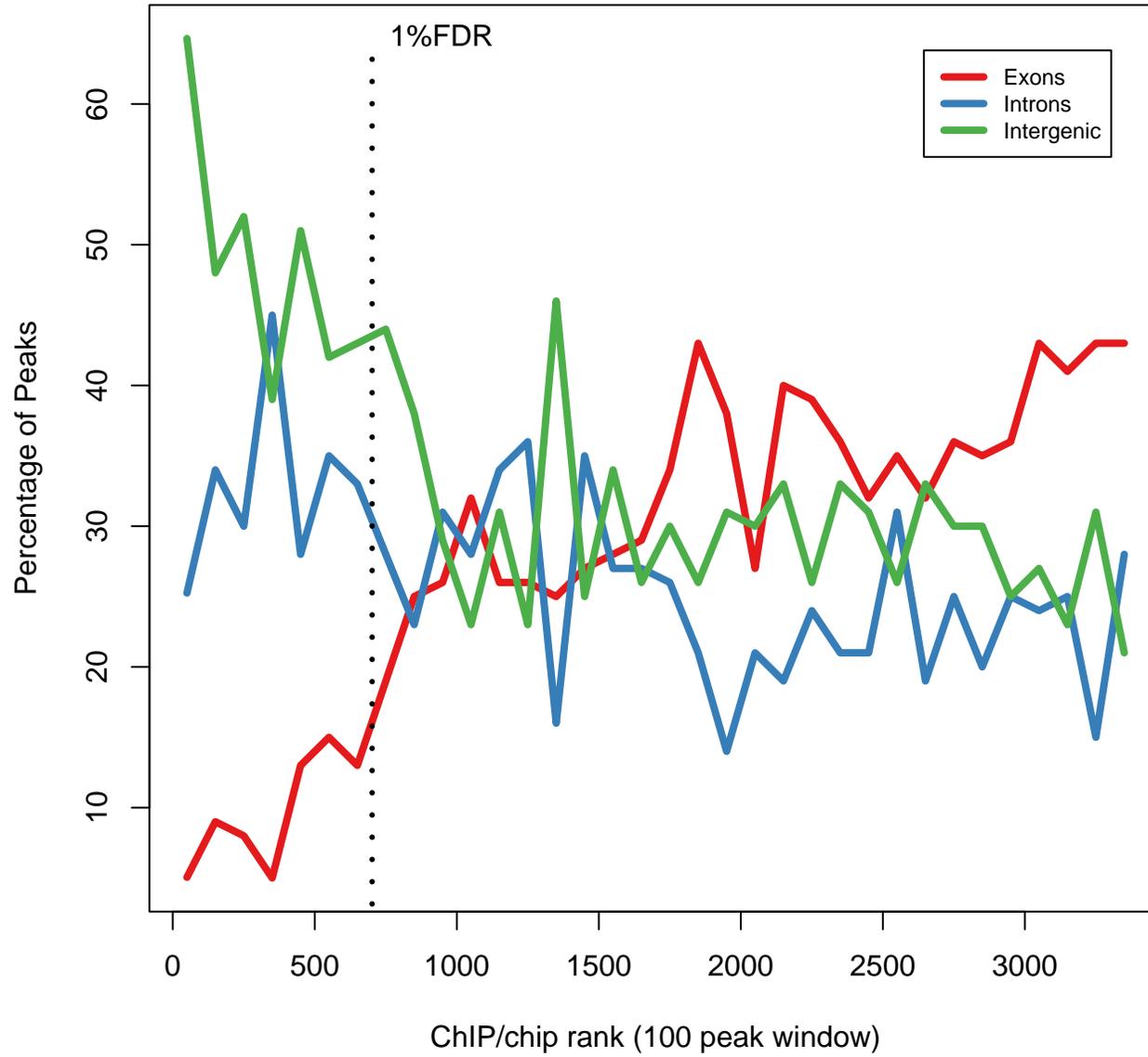
**bcd\_2 PWM enrichment across peaks**



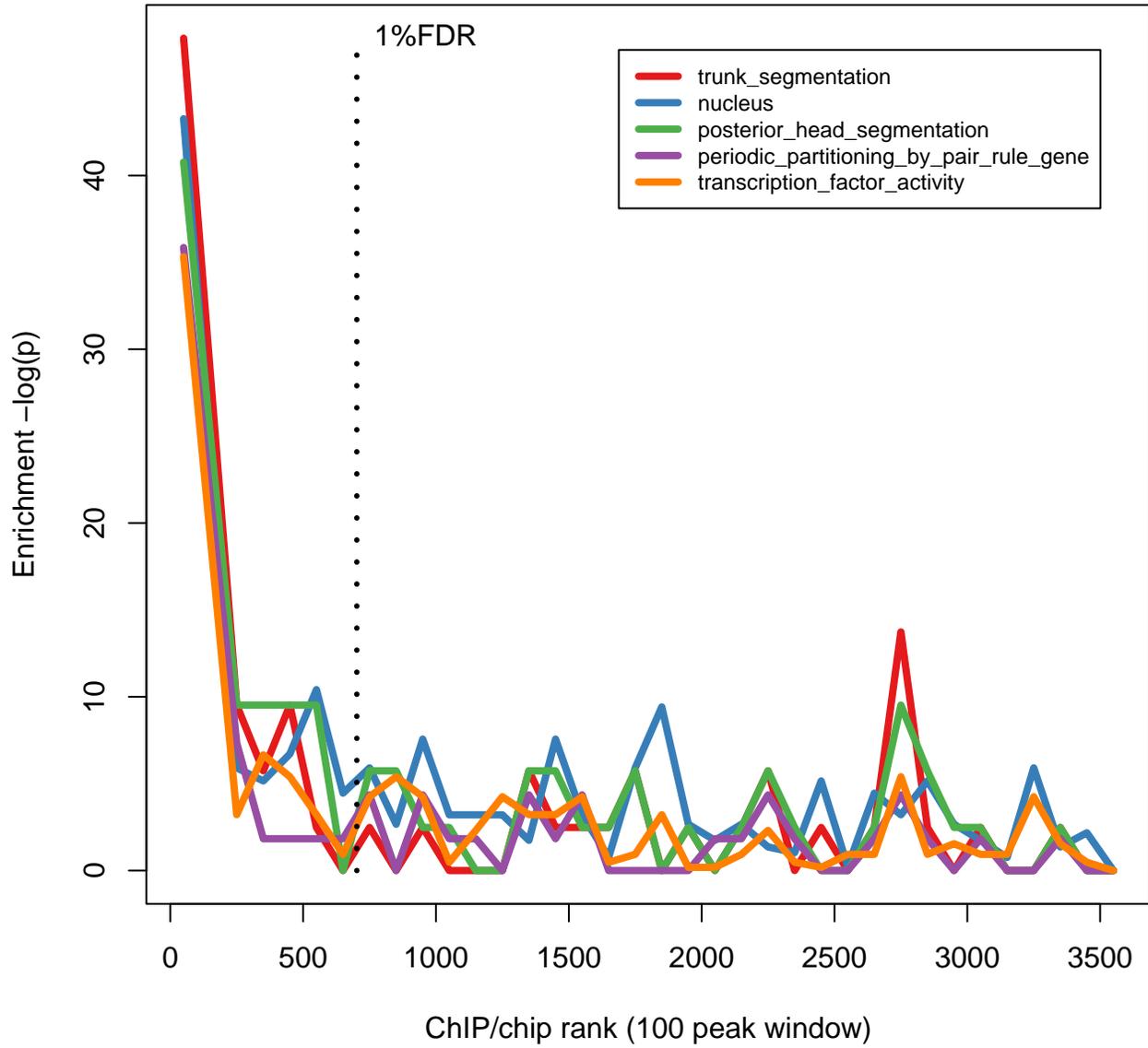
### bcd\_2 PWM enrichment down ranks

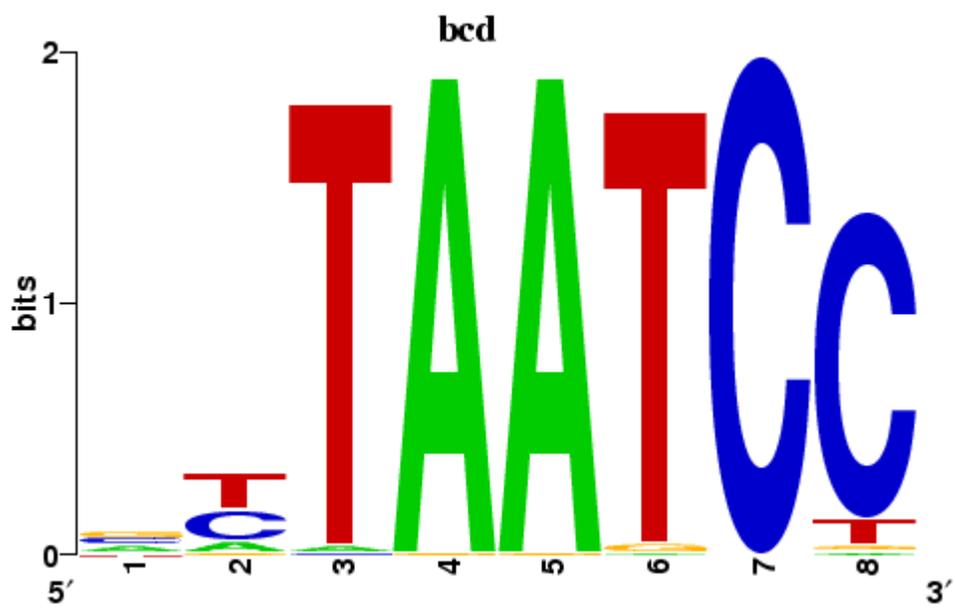


### bcd\_2 genomic location

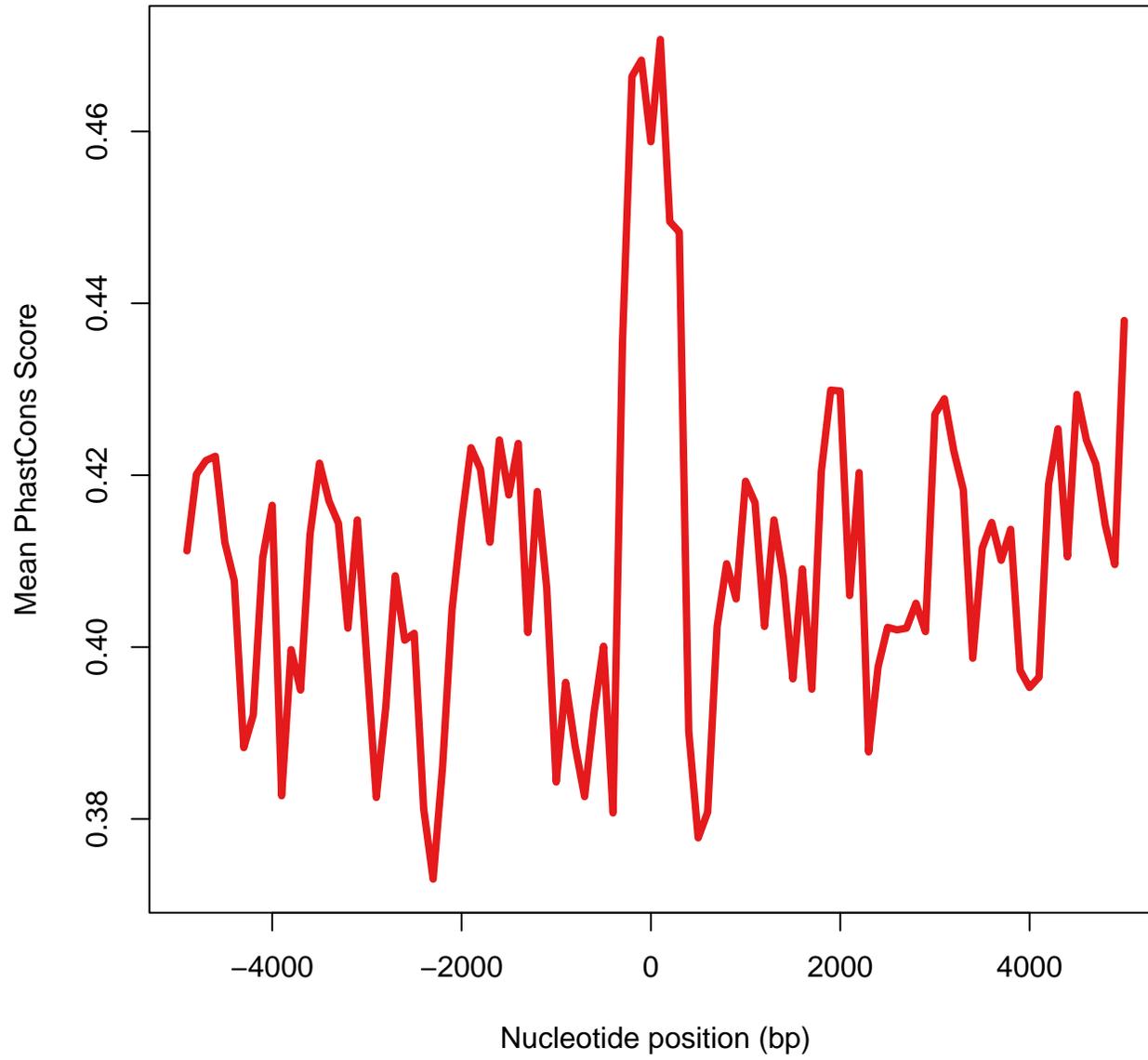


### bcd\_2 GO term enrichment (-logP)





**bcd\_2 mean PhastCons across peaks**



**bcd\_2 mean PhastCons down ranks**

